

**Amendments to the Sequence Listing**

Please replace the originally filed sequence listing with the amended sequence listing attached to this amendment. All amendments relate to improper numbering or improper naming of the modifications to the sequences. Thus, no new matter is added through any of these amendments.

The following amendments were made to the sequence listing:

Seq No.	Originally read	As amended reads	Reason for amendment; basis in specification
26	<220> <221> MOD_RES <222> (5)..(5) <223> AMIDATION	<220> <221> MOD_RES <222> (8)..(8) <223> AMIDATION	misnumbered; see Table 1, Compound 26
65	<220> <221> MOD_RES <222> (1)..(1) <223> N-(2-naphylenesulfonylaminomethyl-4-oxo-butyl) substituted, D	<220> <221> MOD_RES <222> (1)..(1) <223> N-(2-naphylenesulfonylaminomethyl-4-oxo-butyl) substituted	end of phrase ("D form") omitted; divided into two entries; see Table 1, Compound 65
65		<220> <221> MOD_RES <222> (1)..(1) <223> D form	end of phrase ("D form") omitted; divided into two entries; see Table 1, Compound 65
78	<220> <221> MOD_RES <222> (9)..(9) <223> Reduced from amino acid to amino alcohol	<220> <221> MISC_FEATURE <222> (9)..(9) <223> Xaa = Cys reduced from amino acid to amino alcohol	omitted identity of Xaa residue that was reduced; see Table 1, Compound 78
94		<220> <221> DISULFID <222> (3)..(9)	omitted disulfide description; see Table 1, Compound 94
116	<220> <221> MOD_RES <222> (10)..(10) <223> reduced from amino acid to amino alcohol	<220> <221> MISC_FEATURE <222> (10)..(10) <223> Xaa = Ser reduced from amino acid to amino alcohol	omitted identity of Xaa residue that was reduced; see Table 1, Compound 116

Seq No.	Originally read	As amended reads	Reason for amendment; basis in specification
120	<220> <221> MOD_RES <222> (11)..(11) <223> reduced from amino acid to amino alcohol	<220> <221> MISC_FEATURE <222> (11)..(11) <223> Xaa = Pro reduced from amino acid to amino alcohol	omitted identity of Xaa residue that was reduced; see Table 1, Compound 120
122	<220> <221> MOD_RES <222> (11)..(11) <223> reduced from amino acid to amino alcohol	<220> <221> MISC_FEATURE <222> (11)..(11) <223> Xaa = Pro reduced from amino acid to amino alcohol	omitted identity of Xaa residue that was reduced; see Table 1, Compound 122
159	<220> <221> MOD_RES <222> (7)..(7) <223> AMIDATION		incorrectly listed in Table 1 as amidated; should be acid form
166	<220> <221> DISULFID <222> (3)..(8)	<220> <221> DISULFID <222> (3)..(9)	misnumbered; see Table 1, Compound 166
166	<220> <221> MOD_RES <222> (5)..(5) <223> D form	<220> <221> MOD_RES <222> (6)..(6) <223> D form	misnumbered; see Table 1, Compound 166
166	<220> <221> MOD_RES <222> (8)..(8) <223> AMIDATION	<220> <221> MOD_RES <222> (9)..(9) <223> AMIDATION	misnumbered; see Table 1, Compound 166
182	<220> <221> MOD_RES <222> (3)..(3) <223> 4-fluoro substituted, D form	<220> <221> MOD_RES <222> (3)..(3) <223> 4-chloro substituted, D form	incorrectly listed as 4-fluoro rather than 4-chloro; see Table 1, Compound 182